

RAW SEQUENCE LISTINGPATENT APPLICATION: US/09/697,863

DATE: 07/17/2002

TIME: 12:12:09

Input Set : N:\AMC\i697863.raw

Output Set: N:\CRF3\07172002\1697863.raw

Does Not Comply
Corrected Diskette Needed

- 1 <110> APPLICANT: VLAAMS INTERUNIVERSITAIR INSTITUUT VOOR BIOTECHNOL
- 2 <120> TITLE OF INVENTION: CD40-INTERACTING AND TRAF-INTERACTING PROTEINS
- 3 <130> FILE REFERENCE: V7/002-V018
- 4 <140> CURRENT APPLICATION NUMBER: US/09/697,863
- 5 <141> CURRENT FILING DATE: 2000-10-27
- 6 <150> PRIOR APPLICATION NUMBER: 98201392.2
- 7 <151> PRIOR FILING DATE: 1998-04-29
- 8 <160> NUMBER OF SEQ ID NOS: 6
- 9 <170> SOFTWARE: PatentIn Ver. 2.1

ERRORED SEQUENCES

391	<210>	SEQ ID NO: 6															
392	<211>	LENGTH: 442 TYPE: PRT ORGANISM: Homo sapiens															
393	<212>	TYPE: PRT															
394	<213>	ORGANISM: Homo sapiens $/\mathcal{D}$															
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396		Met	Gly	Asp	Pro	Gly	Ser	Glu	Ile	Ile	Glu	Ser	Val	Pro	Pro	Ala	Gly
397		1				5					10					15	
398		Pro	Glu	Ala	Ser	Glu	Ser	Thr	Thr	Asp	Glu	Asn	Glu	Asp	Asp	Ile	Gln
399					20					25					30		
400		Phe	Val	Ser	Glu	Gly	Pro	Ser	Arg	Pro	Val	Leu	Glu	Tyr	Ile	Asp	Leu
401				35					40					45			
402		Val	Cys	Gly	Asp	Asp	Glu	Asn	Pro	Ser	Ala	Tyr	Tyr	Ser	Asp	Ile	Leu
403			50					55					60				
404		Phe	Pro	Lys	Met	Pro	Lys	Arg	Gln	Gly	Asp	Phe	Leu	His	Phe	Leu	Asn
405		65					70					75					80
406		Met	Lys	Lys	Val	Lys	Thr	Asp	Thr	Glu	Asn	Asn	Glu	Val	Ser	Lys	Asn
407						85					90					95	
408		His	Cys	Arg	Leu	Ser	Lys	Ala	Lys	Glu	Pro	His	Phe	Glu	\mathtt{Tyr}	Ile	Glu
409					100					105					110		
410		Gln	Pro	Ile	Ile	Glu	Glu	Lys	Pro	Ser	Leu	Ser	Ser	Lys	Lys	Glu	Ile
411				115					120					125			
412		Asp	Asn	Leu	Val	Leu	Pro	Asp	Cys	${\tt Trp}$	Asn	Glu	Lys	Gln	Ala	Phe	Met
413			130					135					140				
414		Phe	Thr	Glu	Gln	Tyr	Lys	Trp	Leu	Glu	Ile	Lys	Glu	Gly	Lys	Leu	Gly
415		145					150					155					160
416		Cys	Lys	Asp	Cys	Ser	Ala	Val	Arg	His	Leu	Gly	Ser	Lys	Ala	Glu	Lys
417						165					170					175	
418		His	Val	His	Val	Ser	Lys	Glu	Trp	Ile	Ala	Tyr	Leu	Val	Thr	Pro	Asn
419					180					185					190		
420		Gly	Ser	Asn	Lys	Thr	Thr	Arg	Gln	Ala	Ser	Leu	Arg	Lys	Lys	Ile	Arg

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	421			195					200					205						
	422	Glu	His	Asp	Val	Ser	Lys	Ala	His	Gly	Lys	Ile	Gln	Asp	Leu	Leu	Lys			
	423		210					215					220							
	424	Glu	Ser	Thr	Asn	Asp	Ser	Ile	Cys	Asn	Leu	Val	His	Lys	Gln	Asn	Asn			
	425	225					230					235					240			
	426	Lys	Asn	Ile	Asp	Ala	Thr	Val	Lys	Val	Phe	Asn	Thr	Val	Tyr	Ser	Leu			
	427					245					250					255				
	428	Val	Lys	Hís		Arg	Pro	Leu	Ser		Ile	Glu	Gly	Ala		Glu	Leu			
	429	_	- -		260	_		_		265			- •		270		_			
	430	Gln	Glu	_	Asn	Gly	Glu	Val	Asn	Cys	Leu	Asn	Thr	_	Tyr	Ser	Ala			
	431		_	275					280	_			_	285						
	432	Thr		Ile	Ala	Glu	His		Ala	Lys	Glu	Met		Met	Lys	Ile	Phe			
	433	_	290	-1	1	~ 3		295		_	~ 7 .	_	300	- - 7 -	-1		~ 1			
	434	_	Asn	ITe	lle	GLu		Asn	Ala	Lys	He	_	ITe	TTe	TTE	Asp				
	435	305	a	m la	77 - 7	a	310	T	шу	m la	T	315	-1 -	m	T	01	320			
	436	Ala	ser	THY	val		ьуs	ьуs	Thr	Thr	_	val	TTE	TYL	Leu		Cys			
	437	mh x	т1 о	Cln	Cor	325	Dro	ח 7 ח	Dro	17-1	330	Tou	Dho	77 n 1	ת 7 ת	335	Tara			
	438 439	J 11T	116	GTII	340	Ата	PIO	нта	Pro	345	Met	Leu	Pile	vaı	350	ьеu	гур			
	440	Glu	Lou	Wa l		Thr	Tla	71 a	Glu		Tla	Val	Aen	Thr		T.AII	Thr			
	441	Giu	Leu	355	Ser	1111	116	Ата	360	Cys	116	Val	ASII	365	цец	пеа	T 11T			
	442	ጥክ r	T.011		Δen	Cvs	Glv	Dho	Thr	Δan	Glu	Пvr	T. 2 11		Δla	Δsn	T.611			
	443	1111	370	21411	1155	010		375	- 111	11011	OLU	_	380	210	1114	11011	Dou			
E>		Tle		Phe	Cvs	Ser	Asp		Ala	Asn	Thr		i	Glv	Ara	Lvs	Ser			
_ ,	445	385			-1-	502	390	V -1		4121		395		4-1	9	_10	400			
	446		Val	Ala	Thr	Lys		Leu	Glu	Asn	Phe	Pro	Glu	Ile	Ile	Ile	Trp			
	447	-				405					410					415	•			
	448	Asn	Cys	Leu	Asn	His	Arg	Leu	Gln	Leu	Ser	Leu	Asp	Asp	Ser	Ile	Ser			
	449		-		420					4/25	.)				430					
E>	450	Glu	Ile	Lys	Gln	Ile	Asn	His	Leu	Xaa	Tyr									
	451			435					440		/			n					_	. 💉
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																		-		

VARIABLE LOCATION SUMMARY

PATENT APPLICATION: US/09/697,863 TIME: 12:12:10

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Use of n's or Xaa's (NEW RULES):

Use of n's and/or Xaa's have been detected in the Sequence Listing.
Use of <220> to <223> is MANDATORY if n's or Xaa's are present.
in <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

Seq#:1; N Pos. 1627,1849

6 k

Seq#:5; N Pos. 1392,1530,1531

Seq#:5; Xaa Pos. 395,441
Seq#:6; Xaa Pos. 395,441